SEQUENCE LISTING LEO <110> FREN VAN DER LOGABE CORNELIS P. <120> METHOD FOR PRODUCING ANTIBODY FRAGMENTS <130> 060113-0271592 <140> 09/626,242 <141> 2000-09-27 <150> PCT/EP99/00481 <151> 1999-01-25 <150> EP 98300525.7 <151> 1998-01-26 <160> 18 <170> PatentIn Ver. 2.1 <210> 1 <211> 22 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Primer <400> 1 aggtsmarct gcagsagtcw gg 22 <210> 2 <211> 57 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Primer <400> 2 catgccatga ctcgcggccc agccggccat ggccsaggts marctgcags agtcwgg 57 <210> 3 <211> 53

<212> DNA

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<211> 53
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<211> 117
<212> PRT
<213> Lama glama
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Phe Leu Arg Phe Ser Cys Ala Ala Leu Gly Ala Arg Phe Ser Ser Asp
Val Met Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val
Ala Ala Ser Ser Trp Asn Gly Asp Thr Thr His Tyr Ser Asp Ser Val
    50
Glu Gly Gln Phe Thr Ile Ser Arg Asp Ile Ala Lys Asn Thr Ser Tyr
Leu Gln Met Asn Arg Leu Gln Pro Glu Asp Thr Ala Val Tyr Tyr Cys
Arg Trp Cys Arg Pro Pro Arg Pro Lys Tyr Trp Gly Gln Gly Thr Gln
Val Thr Val Ser Ser
        115
<210> 6
<211> 115
<212> PRT
<213> Lama glama
Gln Val Gln Leu Gln Gln Ser Gly Gly Gly Leu Val Gln Ala Gly Ser
Phe Leu Ser Phe Ser Cys Thr Ala Ser Gly Arg Thr Phe Ser Asn Tyr
Ala Met Gly Trp Phe Arg Gln Ala Ser Gly Asn Gln Arg Ala Phe Val
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35

Ala Ala Ile Gly Arg Asn Gly Asp Thr His Tyr Ile Asp Ser Val Lys 50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Gly Lys Asp Thr Val Tyr Leu 65 70 75 80

Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Arg 85 90 95

Ile Trp Val Gly Ala Arg Asp Tyr Trp Gly Gln Gly Thr Gln Val Thr
100 105 110

Val Ser Ser 115

<210> 7

<211> 116

<212> PRT

<213> Lama glama

<400> 7

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly

1 5 10 15

Phe Leu Arg Phe Ser Cys Ala Ala Ser Gly Arg Thr Phe Ser Arg Tyr

Thr Met Gly Trp Phe Arg Gln Ala Pro Gly Asn Glu Arg Lys Phe Val 35 40 45

Ala Ala Val Ser Thr Ser Gly Asn Thr His Tyr Thr Gly Ser Val Lys
50 55 60

Gly Arg Phe Thr Ile Phe Arg Gln Asn Ala Lys Asn Thr Val Tyr Leu 65 70 75 80

Gln Met Ser Asn Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Ala 85 90 95

Ala Arg Phe Gly Gly Met Asn Trp Lys Tyr Trp Gly Gln Gly Ile Gln 100 105 110

Val Thr Val Ser 115

<210> 8

<211> 121

<212> PRT

<213> Lama glama

<400> 8

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Pro

1 5 10 15

Phe Leu Asn Val Ser Cys Val Val Ser Gly Gly Ile Phe Ser Asp Tyr
20 25 30

01

Thr Leu Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Lys Phe Val

Ala Ala Val Ser Ser Gly Gly Ser Thr His Tyr Thr Gly Ser Val Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Ala Asn Thr Met Tyr Leu 65 70 75 80

Gln Met Ser Ser Leu Lys Pro Asp Asp Thr Ala Val Tyr Tyr Cys Asn 85 90 95

Ala Ile Val Pro Pro Thr Arg Thr Phe Cys Gly Arg Thr Tyr Trp Gly
100 105 110

Gln Gly Thr Gln Val Thr Val Ser Ser 115 120

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<212> PRT

<213> Lama glama

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Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Asp 1 5 10 15

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Ala Val Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val

Gly Arg Ile His Arg Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Thr Gln Asn Thr Val Tyr 65 70 75 80

Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Asn Val Arg Ser Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser . 100 105 110

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<211> 117

<212> PRT

<213> Lama glama

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Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly
1 5 10 15

Phe Leu Arg Phe Ser Cys Ala Ala Ser Asn Ala Leu Phe Ser Gly Tyr 20 25 30

Ala Met Gly Cys Phe Arg Gln Ala Val Gly Lys Glu Arg Glu Phe Val 35 40 45

Ala Ala Ile Thr Trp Asn Asn Arg Asn Thr His Tyr Ala Asp Ser Val 50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Val Tyr 65 70 75 80

Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys 85 90 95

Thr Ser Gly Met Arg Arg Leu Gly Asp Tyr Trp Gly Gln Gly Thr Gln
100 105 110

Val Thr Val Ser Ser 115

<210> 11

<211> 124

<212> PRT

<213> Lama glama

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Gln Val Lys Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Asp Lys Tyr
20 25 30

Ala Ile Gly Trp Phe Arg Gln Ala Pro Gly Lys Gln Arg Glu Leu Val 35 40 45

Ala Gly Ile Ser Thr Gly Gly Ser Thr Asn Tyr Ala Asp Ser Val Lys
50 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asp Thr Val Tyr Leu 65 70 75 80

Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Ala 85 90 95

Ala Gly Arg Arg Ile Ser Ser Ser Tyr Tyr Ser Arg Gly Leu Tyr Ala 100 105 110

Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser 115 120

<210> 12

<211> 124

<212> PRT

<213> Lama glama

01

<400> 12

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Asp 1 5 10 15

Ser Leu Arg Leu Ser Cys Glu Ala Ser Gly Arg Ser Phe Ser Asn Phe 20 25 30

Ala Met Ala Trp Phe Arg Gln Thr Pro Gly Lys Glu Arg Glu Phe Val 35 40 45

Ala Gly Ile Ser Trp Arg Gly Gly Arg Thr Tyr Tyr Ala Ala Ser Val
50 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Gly Lys Asn Thr Val Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys 85 90 95

Ala Thr Ala Tyr Gly Gln Gly Pro Ile Thr Val Pro Lys Phe Tyr Thr 100 105 110

Tyr Arg Gly Gln Gly Thr Gln Val Thr Val Ser Ser 115 120

<210> 13

<211> 121

<212> PRT

<213> Lama glama

<400> 13

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly
1 5 10 15

Cys Val Arg Leu Ser Cys Ala Ala Ser Gly Arg Thr Phe Ser Arg Tyr

Thr Met Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val 35 40 45

Ala Ala Ile Ser Trp Arg Ser Gly Gly Ile Lys Ile Tyr Gly Asp Ser 50 60 .

Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asp Thr Val

Tyr Val Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr 85 90 95

Cys Asn Ser Arg Pro Arg Ile Tyr Arg Gly Asn Val Val Tyr Trp Gly
100 105 110

Gln Gly Thr Gln Val Thr Val Ser Ser

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       peptide
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                  5
 <210> 16
 <211> 39
 <212> DNA
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       oligonucleotide
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 gcggccgccc atcaccatca ccatcacggg gccgcagaa
                                                                     39
<210> 17
'<211> 13
 <212> PRT
 <213> Artificial Sequence
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 <223> Description of Artificial Sequence: Synthetic
      peptide
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Ala Ala Ala His His His His His Gly Ala Ala Glu
                   5
<210> 18
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<213> Unknown Organism
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<220>

<223> Description of Unknown Organism: Myc peptide sequence

<400> 18

Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn 1 5 10